INFORMAL SEQUENCE LISTING

SEQ ID NO:1
Wild type DT with signal sequence underlined

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SEQ ID NO:2

30 DT-GML1

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SEQ ID NO:4

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SEQ ID NO:5

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SEQ ID NO:9

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SEQ ID NO:10

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SEQ ID NO:11
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SEQ ID NO:12
DT-IL2U2
ATGGGCGCCGACGACGTCGTCGACTCTTCTAAATCTTTTGTGATGGAAAACTTTTCTTCGTACCACGGGACTAAA
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TGGAAAGGGTTTTATAGTACCGACAATAAATACGACGCTGCGGGATACTCTGTAGATAATGAAAACCCGCTCTCT
GGAAAAGCTGGAGGCGTGGTCAAAGTGACGTATCCAGGACTGACGAAGGTTCTCGCACTAAAAGTGGATAATGCC

GAAACTATTAAGAAAGAGTTAGGTTTAAGTCTCACTGAACCGTTGATGGAGCAAGTCGGAACGGAAGAGTTTATC

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CAAGATGCGATGTATGAGTATATGGCTCAAGCCTGTGCAGGAAATGGAAGTGGAAGATCAGCAGGTAGCTCATTG

TCATGCATAAATCTTGATTGGGATGTCATAAGGGATAAAACTAAGACAAAGATAGAGTCTTTGAAAGAGCATGGC

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TTTCATCAAACGGCATTAGAGCATCCTGAATTGTCAGAACTTAAAACCGTTACTGGGACCAATCCTGTATTCGCT

GGGGCTAACTATGCGGCGTGGGCAGTAAACGTTGCGCAAGTTATCGATAGCGAAACAGCTGATAATTTGGAAAAG

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TTAATCAGCAATATCAACGTAATAGTTCTGGAACTAAAGGGATCTGAAACAACATTCATGTGTGAATATGCTGAT

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SEQ ID NO:13

DT-IL2U3

ATGGGCGCCGACGACGTCGTCGACTCTTCTAAATCTTTTGTGATGGAAAACTTTTCTTCGTACCACGGGACTAAA

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CAAGATGCGATGTATGAGTATATGGCTCAAGCCTGTGCAGGAAATGGAAGTGGAAAATCAGCAGGTAGCTCATTG

TCATGCATAAATCTTGATTGGGATGTCATAAGGGATAAAACTAAGACAAAGATAGAGTCTTTGAAAGAGCATGGC

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TATAATCGTCCCGCGTATTCTCCCGGGCATAAAACGAGGCCTCATATGGCACCTACTTCAAGTTCTACAAAGAAA

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AAACTCACCAGGATGCTCACATTTAAGTTTTACATGCCCAAGAAGGCCACAGAACTGAAACATCTTCAGTGTCTA

GAAGAAGAACTCAAACCTCTGGAGGAAGTGCTAAATTTAGCTCAAAGCAAAAACTTTCACTTAAGACCCAGGGAC

TTAATCAGCAATATCAACGTAATAGTTCTGGAACTAAAGGGATCTGAAACAACATTCATGTGTGAATATGCTGAT

GAGACAGCAACCATTGTAGAATTTCTGAACAGATGGATTACCTTTTGTCAAAGCATCATCTCAACACTGACTTGA

SEQ ID NO:14
T7 promoter primer (5' primer for DT constructs)

5   GTAATACGACTCACTATAGGGC

SEQ ID NO:15
3' mutagenic primer for U2 constructs

10   GATTTATGCATGACAATGAGCTACCTGCTGATCTTCCACTTCCATTTCCTGCACAGGCTTG

SEQ ID NO:16
3' mutagenic primer for U3 constructs

15   GATTTATGCATGACAATGAGCTACCTGCTGATTTTCCACTTCCATTTCCTGCACAGGCTTG

SEQ ID NO:17
3' mutagenic primer for L1 constructs

20   GATTTATGCATGACAATGAGCTACCTTGACTCAACATTCCTAATGGTCCATTTCCTGCACAGGCTTG

SEQ ID NO:18
3' mutagenic primer for L2 constructs

25   GATTTATGCATGACAATGAGCTACCTTGTGCCCATAATCCTAATGGTCCATTTCCTGCACAGGCTTG

SEQ ID NO:19
MMP substrate octapeptide for L1 constructs

30   GPLGMLSQ

SEQ ID NO:20
MMP substrate octapeptide for L2 constructs

35   GPLGLWAQ

SEQ ID NO:21
uPA substrate hexapeptide for U2 constructs

40   GSGRSA

SEQ ID NO:22
uPA substrate hexapeptide for U3 constructs

45   GSGKSA